

PCT

RAW SEQUENCE LISTING DATE: 05/27/2003 PATENT APPLICATION: US/10/069,410 TIME: 13:31:03

Input Set : A:\12795-009US1.txt

Output Set: N:\CRF4\05272003\J069410.raw

4 <110> APPLICANT: Germain , Stephane Edouard Hill, Caroline Susan Howell, Michael Terence 8 <120> TITLE OF INVENTION: POLYPEPTIDES 11 <130> FILE REFERENCE: 12795-009US1 13 <140> CURRENT APPLICATION NUMBER: US 10/069,410 14 <141> CURRENT FILING DATE: 2002-02-25 16 <150> PRIOR APPLICATION NUMBER: PCT/GB00/03265 17 <151> PRIOR FILING DATE: 2000-08-25 19 <150> PRIOR APPLICATION NUMBER: GB 9920000.8 20 <151> PRIOR FILING DATE: 1999-08-25 22 <160> NUMBER OF SEQ ID NOS: 63

24 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

will it ro. 1237 <210> SEQ ID NO: 63 Dees Not Comply 1238 <211> LENGTH: 16 Corrected Diskette Needed 1239 <212> TYPE: PRT 1240 <213> ORGANISM: Artificial Sequence

1242 <220> FEATURE:

1243 <223> OTHER INFORMATION: Synthetically generated peptide

1245 <400> SEQUENCE: 63

1246 Ser Lys Pro Thr Thr Lys Gln Arg Gln Asn Lys Pro Pro Asn Lys Pro
1247 1 5 10 IStimut. E--> 1247 1

VERIFICATION SUMMARY DATE: 05/27/2003 PATENT APPLICATION: US/10/069,410 TIME: 13:31:04

Input Set : A:\12795-009US1.txt

Output Set: N:\CRF4\05272003\J069410.raw

```
L:326 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!

L:330 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24

L:334 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24

L:339 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24

L:344 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24

L:349 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24

L:353 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24

L:357 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24

L:361 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24

L:365 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24

L:366 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0

M:341 Repeated in SeqNo=24

L:824 M:351 W: Sequence data Name/Key Feature Out-of-Range, SEQ ID#:54, CDS LOCATION: (1)...

(1575)

L:1247 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:63
```